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- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 794 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: cDNA

- (vii) IMMEDIATE SOURCE:
(B) CLONE: U-D63

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 52:

GGCACGAGGG ATTATAAAAA GAAACTCGCA CTCAGCCAGA TTGACTCAAG CATGAAGGTT 60
GCATCTTTTCG AATGTTCCCTT TTTTTCCTCAA CATCTTTTGA ATGTTACATG ACAATCATGA 120
CGCCGTGCAT ACAATGATAC CTCTGATAGG GTTGAATGCA CAAACCACTC ATCGTGCTAT 180
ATTGCAAGTG CTTATGTTTG TATTCATTTT CTTATAATTT CAATGTGAGA GTTACCTTAG 240
GTTTCATATTA NATTTATATA ATATTGGGCA TAGATTTTTT TGTCATTGTT TGTGATCTGC 300
AATTTTCATTA GGTCATCGAC CATATTATGG TAAATTTAAT TTACTAAATT ATATCTTGAG 360
TTTATTGCTT TGCATATTTT GTGCTGGAAA CCATGCCAAT CCAACTCCTT TGCAAAACTT 420
AAGATAATTA AAGGAGATAT TGAGGATGCC GACTGCTTTA GGTTATTTGC ATTANACAAT 480
AAACGCCCCGT GAAATGGAAA TCATGCAAAG CTTTGCATGG AGAATATGTT NATAGAGATA 540
GACATGAAGA CATGAATATG CTTTCTTGAT GAACAACGAT GTTACACNTGT TTACTGCATG 600
CATGATACAT CCAATGTTCT AGGCTTGTTG GCAATCATTT TATTCNAAAA ATTGTCTGTT 660
TCTCTNCCNC CATAAGGTTA GCTTGTGGAA AATGTTCAAC TTTGGGCNCA NATGATCCGT 720
TTTAGCAAAA TCCCNCCATT CCTATTTTTT TCCGGAATCC NNTTGGNAAA ANATCCTNAC 780
ATACTATTTT CACA 794

(2) INFORMATION FOR SEQ ID NO: 53:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 798 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: cDNA

- (vii) IMMEDIATE SOURCE:
(B) CLONE: U-D65

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 53:

GGCACGAGGC CCGTAGAGCG AGTGCTGAGA TGGCTTTGAG GGCCTTCTTC CCCCTCTGCA 60
TCGCTCTCGT GGTGACGCTC TCACCTCTGT GTGATGCCAC TTCGCCTTAC TACACCATCA 120
CACCGCCAC CCGCGTGGCC AAGCCGCCTT CAGTTGAACC ACCACCCTAC CACGGCCCTC 180
CGACGACCCA CCCTAAGCCA CCGAGTCATG GTGGCCAACC TCCGTCCAC CATCACCCAA 240

CACCAATCTA CGGTGCACCC CCTCCGCAAC ACCACCACCA CCACCAACAC CACCACCAAC 300
CTGCACCACC AACTCACGCA GAACACCCTC CGTACTACCA CATGCCTTCC CCGCCGCCGC 360
5 ATGGCCAGCA CCCGTCACCA CCGTCACATG ATTATCCCGT ACCTCCTGCT CACAAGCCCC 420
GAACTCCGCC GCCGGTTTAC AAGTCTCCAC CACCGACCCA CCGTCCTTAC CCTCCATCGA 480
CGCCACCCCA CCATCCGACG CACCCGCCTT CTCAGCCGAC GCCGTCATAC AAGGCCCCGC 540
10 CACCATACAA GAACATCCCT GANCATCTCC ACCGCCGCGT CACTATCATT CTCCGTCTTC 600
ACCACCANCA CAACCACCAT NCAATAGTC TCGTTTGCAT CTCTCCGTTG AANATGAACC 660
15 AATGTCNTTT AATAACGATC AGGGTTTCAA ATAAAAACNA ATTTCCGCCA TTGTAATGCT 720
ATGGTTGTTC TCTCTGCTTC CCGGGGAAAG TTTCTTGGGT CATNTTAACC NCCTCCTAAT 780
GTTGNGCTCT TNNTANAA 798

(2) INFORMATION FOR SEQ ID NO: 54:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 824 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: cDNA

(vii) IMMEDIATE SOURCE:
(B) CLONE: U-D84

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 54:

GGCACGAGCG CCCACCCCGG TGGCCAAGCC GCCTTCAGTT GAACCACCAC CCTACCACGG 60
CCCTCCGACG ACCCACCCTA AGCCACCGAG TCATGGTGGC CAACCTCCGT CCCACCATCA 120
CCCAACACCA ATCTACGGTG CACCCCTCC GCAACACCAC CACCACCACC AACACCACCA 180
45 CCAACCTGCA CCACCAACTC ACGCAGAACA CCCTCCGTAC TACCACATGC CTCCCCGCC 240
GCCGCATGGC CAGCACCCGT CACCACCGTC ACATGATTAT CCCGTACCTC CTGCTCACAA 300
GCCCCGAACT CCGCCGCCGG TTTACAAGTC TCCACCACCG ACCCACCGTC CTTACCCTCC 360
50 ATCGACGCCA CCCACCATC CGACGCACCC GCCTTCTCAG CCGACGCCGT CATAAAGGC 420
CCCGCCANCA TNCNAGACCA TCCCTTGAGC ACTTCTCCAN CGNTTGC GTT TATTATCATT 480
55 CTCCGTCTTC ACCACCACCA CCACCACCAT ACAAATAGTC TCGTTTGCCA TCTCTCCGTT 540
GAAGATGACG CAGTGTGCTT TTAGTAGCGA TCAGGGTTAC CAATAAGAAC GATGTTGCCG 600
CCATTGTAAT GGCTATGGTT GTTCTCTCTG CTTTTCCGGG GGAAGGTTCT TTGGGTTTCA 660
60 GTTAAACCTC TCTCTTAAAT GTTCATGCAT CTTATTATAA ACNAAAATTG GCCATTTNNN 720
NNNTNTNNTN NNNNNTTNNN NNNNNAAAAC TCNAAAATA TTTTAAAAA CGGGCGGGGG 780
65 GCCCATCNAT TTTCCNNCCC GGGTNGGGGN TCCAGNTTA TTNT 824

(2) INFORMATION FOR SEQ ID NO: 55:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 797 base pairs
(B) TYPE: nucleic acid

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(C) STRANDEDNESS: single
(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: cDNA

(vii) IMMEDIATE SOURCE:
(B) CLONE: U-D75

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 55:

GGCACGAGGG	TTACGATCGT	CATGTTGGGG	GTGTTACGCG	GGGNAGGTGG	TGGAGGTGCC	60
GGCGGAGCTG	GTGGCCGCCG	GCAGCAGGAC	GCCGTCCCTT	AAGACACGAG	CGTCGGAGCT	120
GATCAACCGA	TTCCTCGGGA	GCTTCGCTCC	CGCGGTGTCT	ATCCAGATCG	GGGACTTGGG	180
ACACCTCGCC	TACTCCACG	CCAACCAGTC	CCCCTTCGCT	CCCAGGTTGT	TTGCAGCGAA	240
GGACGANATT	TACTGCCTCT	TCAAGGGAGT	GCTGACCAAC	CTGGGCAGCT	TGAGGCAGCA	300
GSTATGGGCTT	TCCAAGAGTG	CCGACNAGGT	GGTGTGGTCT	ATCGAAGCCT	ACAAGGCCCT	360
CCGTGACCGA	GCTCCCTATC	CTCCCAGCTT	CATGCTCGCA	CACCTTACTG	GCAACTTCGC	420
CTTCGTGCTC	TTCGACAAGT	CCACATCATC	CATCCTTGTT	GCATCTGACC	CANATGGAAN	480
ANTACCCTTG	TTCTGGGGGA	TACTGCANA	TGGATGCCTT	GCCTTTGCTG	ACNATCTANA	540
CTTGCTGAAG	GGATCGTGCG	GGAAGTCACT	TGCACCATTG	CCTGAAGGAT	GTTACTATTC	600
CAATGCCTTG	GGGGGCCTGA	AAANCTATGA	AAACCCCAAC	ACAAGGTGAA	CTGCTGTTTC	660
TTGAANATGA	AGAANAATTN	TTTGTGCCNC	TTTCAAGGTG	GAANGATCTG	CCNTTCTTGC	720
GGNAACCCCC	TAATCAAGGA	ACATCTTCCA	ATGTTTGGGG	CNAATGATCC	TCTTAAAATC	780
CTAAAATTCT	TGGAATT					797

(2) INFORMATION FOR SEQ ID NO: 56:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 804 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: cDNA

(vii) IMMEDIATE SOURCE:
(B) CLONE: U-D83

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 56:

GGCACGAGCT	TCCACACAGG	TGAGGGTAAG	TCGGTTACGA	TCGTCATGTT	GGGGGTGTTC	60
AGCGGGGAGG	TGGTGGAGGT	GCCGGCGGAG	CTGGTGGCCG	CCGGCAGCAG	GACGCCGTCC	120
CCTAAGACAC	GGGCGTCGGA	GCTGATCAAC	CGATTCTCTG	GGAGCTTCGC	TCCC GCGGTG	180
TCGATCCAGA	TCGGGGACTT	GGGACACCTC	GCCTACTCCC	ACGCCAACCA	GTCCCCCTTC	240
GCTCCCAGGT	TGTTTGAGC	GAAGGACGAN	ATTTACTGCC	TCTTCAAGGG	AGTGCTGACC	300
AACCTGGGCA	GCTTGAGGCA	GCAGTATGGG	CTTTCCAAGA	GTGCCGACNA	GGTGGTGCTG	360